

Serial Number: 09/991,452

CRF Processing Date: 12/3+5/01  
 Edited by: DC  
 Verified by: DC (STIC sta

112

**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

OIPE

## RAW SEQUENCE LISTING

DATE: 12/03/2001

PATENT APPLICATION: US/09/991,452

TIME: 11:06:13

Input Set : A:\#357703 v1 - A34535-A-PCT-USA-A Sequence Listing.txt

Output Set: N:\CRF3\11212001\I991452.raw

**Does Not Comply  
Corrected Diskette Needed**

4 <110> APPLICANT: Fisher, Paul B.  
6 <120> TITLE OF INVENTION: USE OF A MELANOMA DIFFERENTIATION  
7 ASSOCIATED GENE (mda-7) FOR REVERSING A CANCEROUS PHENOTYPE  
10 <130> FILE REFERENCE: 34535-A-PCT-USA-A  
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/991,452  
13 <141> CURRENT FILING DATE: 2001-11-20  
15 <150> PRIOR APPLICATION NUMBER: US 09/251,124  
16 <151> PRIOR FILING DATE: 1999-02-16  
18 <150> PRIOR APPLICATION NUMBER: PCT/US97/14548  
19 <151> PRIOR FILING DATE: 1997-08-15  
21 <150> PRIOR APPLICATION NUMBER: US 08/696,573  
22 <151> PRIOR FILING DATE: 1996-08-16  
24 <160> NUMBER OF SEQ ID NOS: 1  
26 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
28 <210> SEQ ID NO: 1  
29 <211> LENGTH: 12  
30 <212> TYPE: PRT  
31 <213> ORGANISM: Homo Sapiens  
33 <400> SEQUENCE: 1  
34 Pro Ser Gln Glu Asn Glu Met Phe Ser Ile Arg Asp  
35 1 5 10  
37 NY02:357703.1

—delete

VERIFICATION SUMMARY

DATE: 12/03/2001

PATENT APPLICATION: US/09/991,452

TIME: 11:06:14

Input Set : A:\#357703 v1 - A34535-A-PCT-USA-A Sequence Listing.txt

Output Set: N:\CRF3\11212001\I991452.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number